

**Table S1.** Determining corrole:protein ratios of assembled HerMn preparations.\*

<b>S2Mn</b> <sup>†</sup>		<b>HerPBK10</b> <sup>‡</sup>		<b>HerMn</b>	
A ( $\lambda_{\text{max}}$ ) <sup>§</sup>	Conc ( $\mu\text{M}$ )	A (595 nm)	Conc ( $\mu\text{g/mL}$ )	Conc ( $\mu\text{M}$ )	Corrole:Protein Molar Ratio <sup>¶</sup>
0.0246	117.14	0.1610	475.77	4.90	23.9
0.0079	37.62	0.0380	112.29	1.16	32.5
0.1067	508.10	0.4590	1356.38	13.98	36.3

\* HerMn assemblies were analyzed by absorbance spectroscopy to determine the S2Mn and HerPBK10 concentrations in each HerMn preparation. Table shows three typical yields from more than forty independent preparations.

<sup>†</sup> S2Mn concentration was determined by the equation:

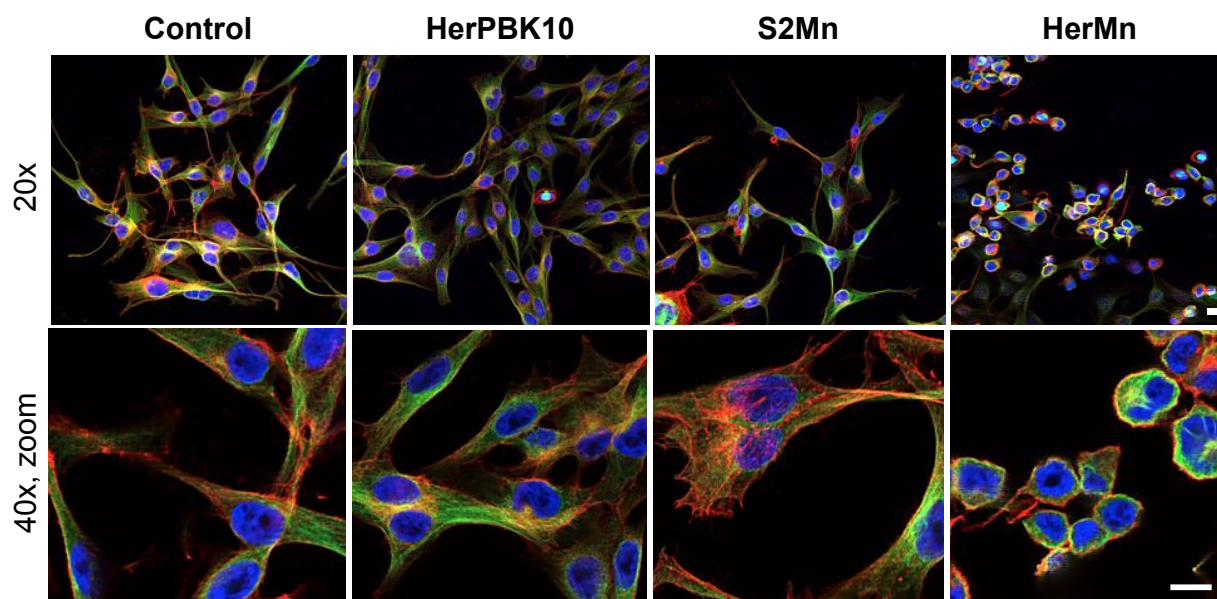
$$[A(\lambda_{\text{max}}) / \epsilon] \times \text{dilution factor} = M \text{ (molar)}$$

where  $\epsilon$  (extinction coefficient) is  $21,000 \text{ M}^{-1}\text{cm}^{-1}$  when absorbance is measured using a 1 cm path-length cuvette.

<sup>‡</sup> HerPBK10 concentration was determined by Bio-Rad (Bradford-based) Protein Assay, at A595 and extrapolated against a standard calibration curve.

<sup>§</sup>  $\lambda_{\text{max}} = 420 \text{ nm}$ . An absorbance spectrum was taken of each HerMn preparation to determine the  $\lambda_{\text{max}}$ , which consistently peaked at 420 nm.

<sup>¶</sup> Molar Ratio = # of corrole molecules bound per protein molecule.



**Supplemental Figure S1. Effect of HerMn on the cytoskeleton.** Low (20x) and higher (enlargement of 40x region) magnification of HER2+ MDA-MB-435 cells after treatment for 24 hours with HerMn or S2Mn (each at 5  $\mu$ M corrole concentration), HerPBK10 (at equivalent protein concentration as HerMn) or PBS (control), followed by washing, fixation, and processing for immunofluorescence. Images were obtained using a Leica SPE laser scanning confocal microscope. Red, actin; Green, tubulin; Blue, nucleus. Scale bar= $\sim$ 10 microns.